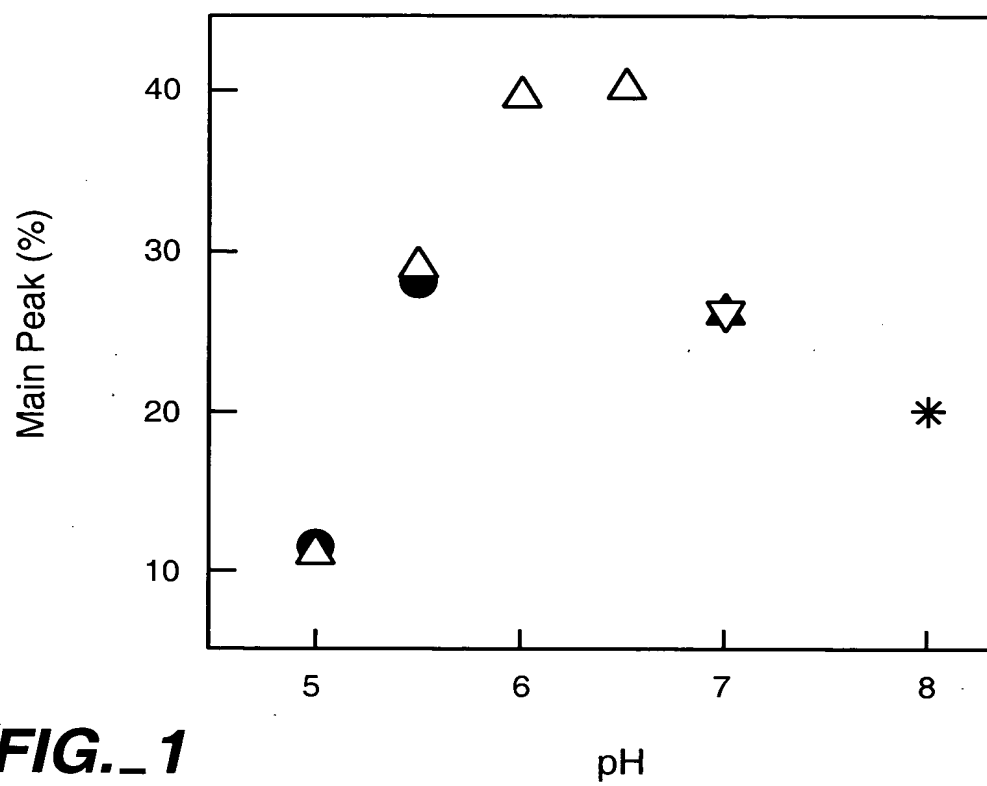
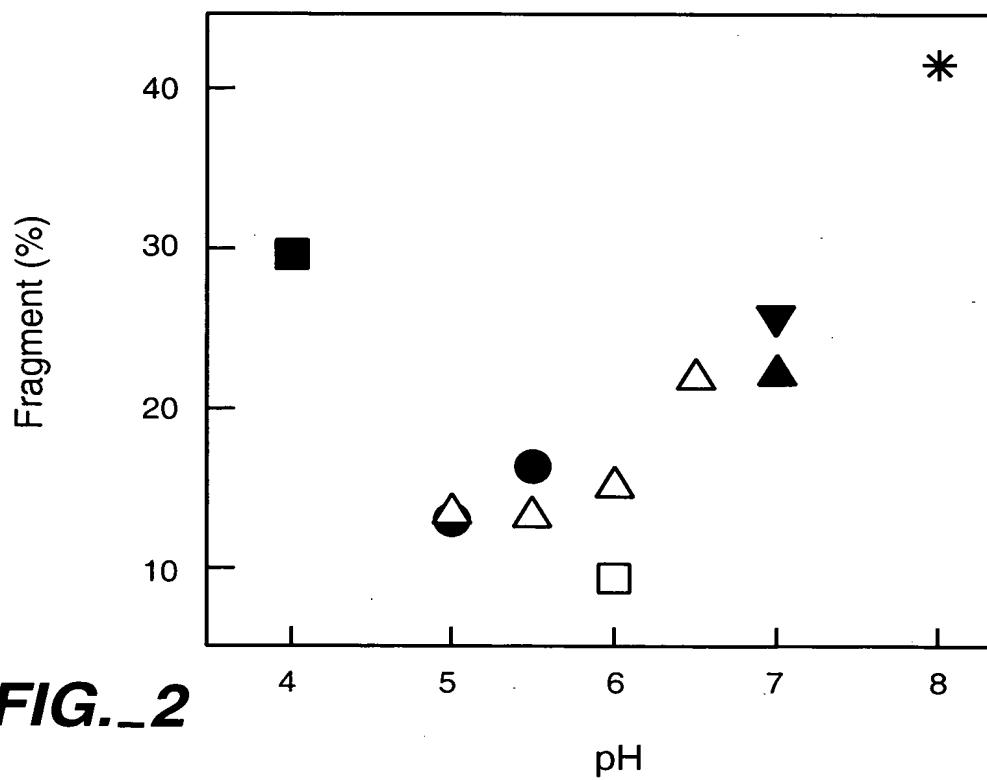
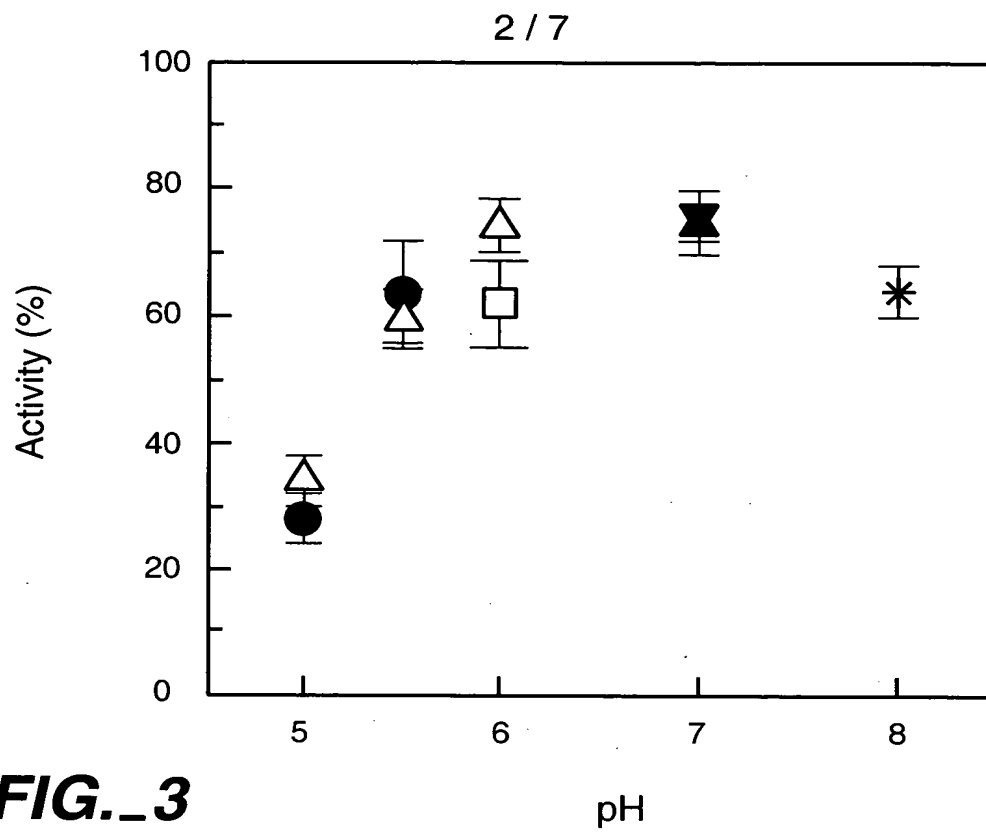
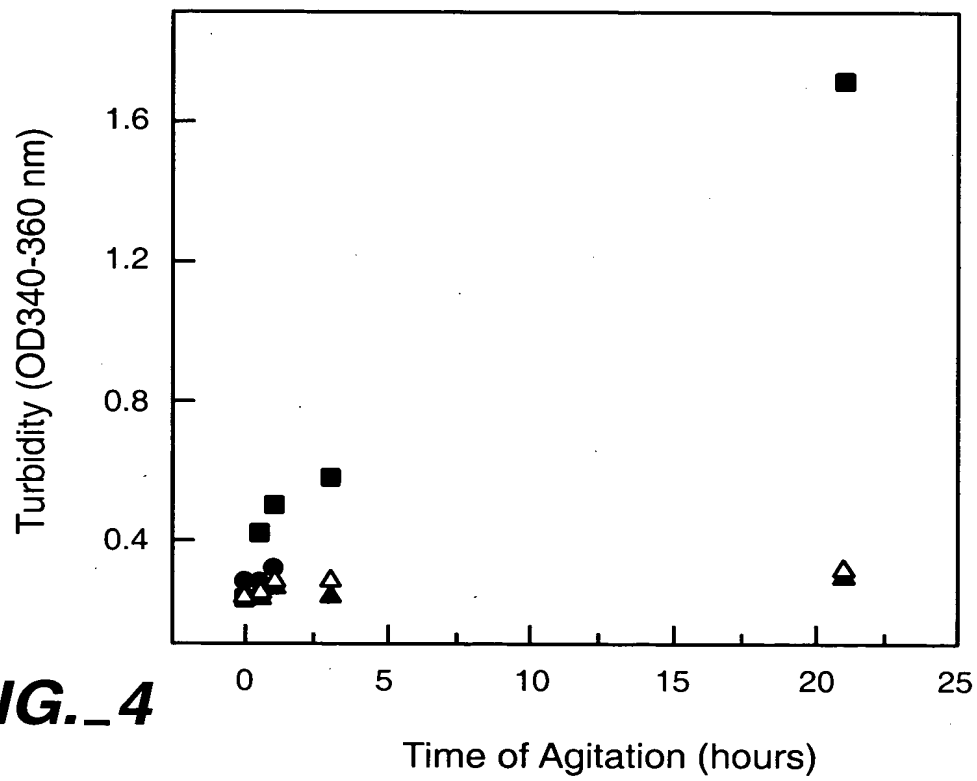
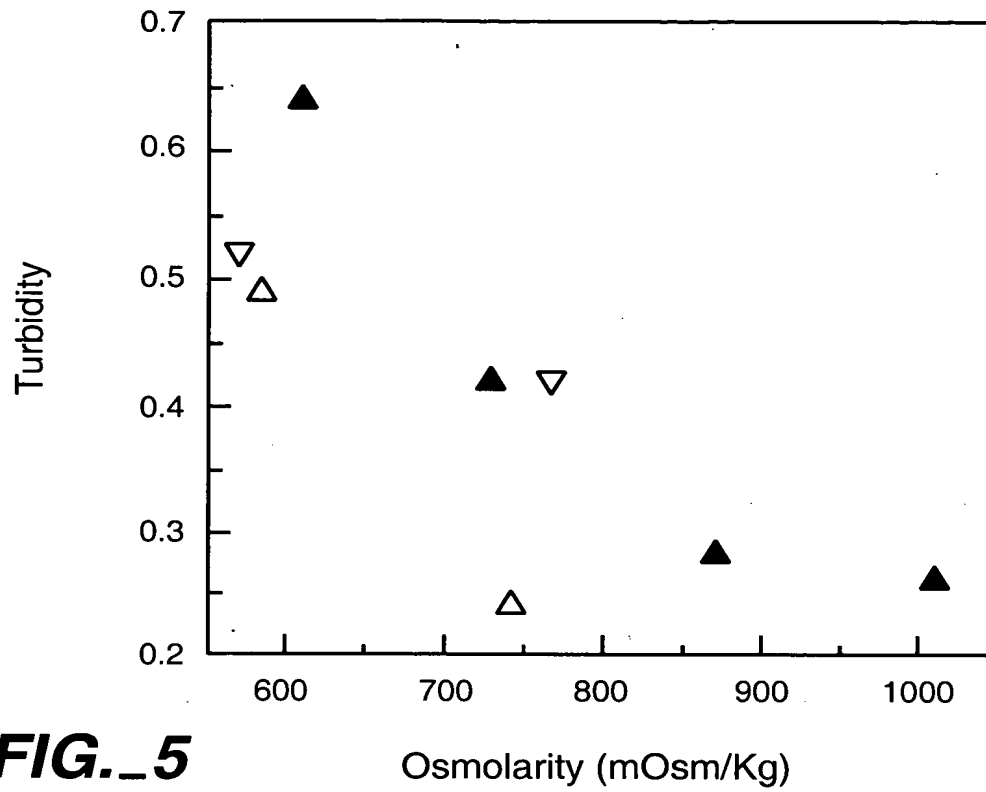
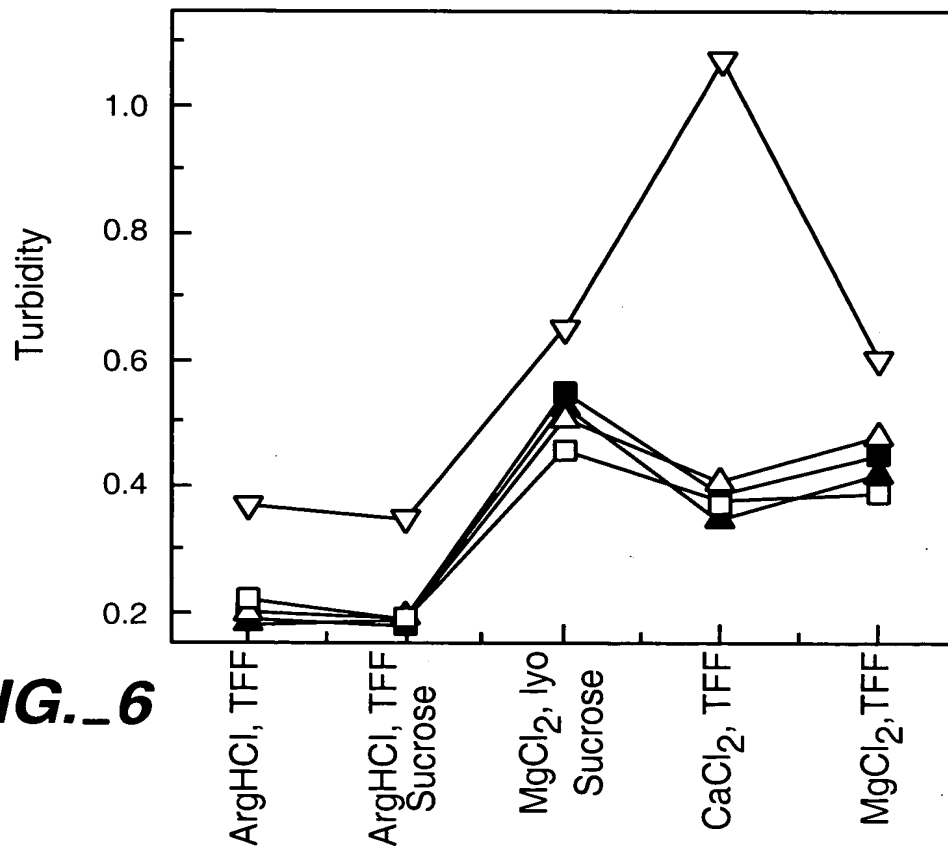


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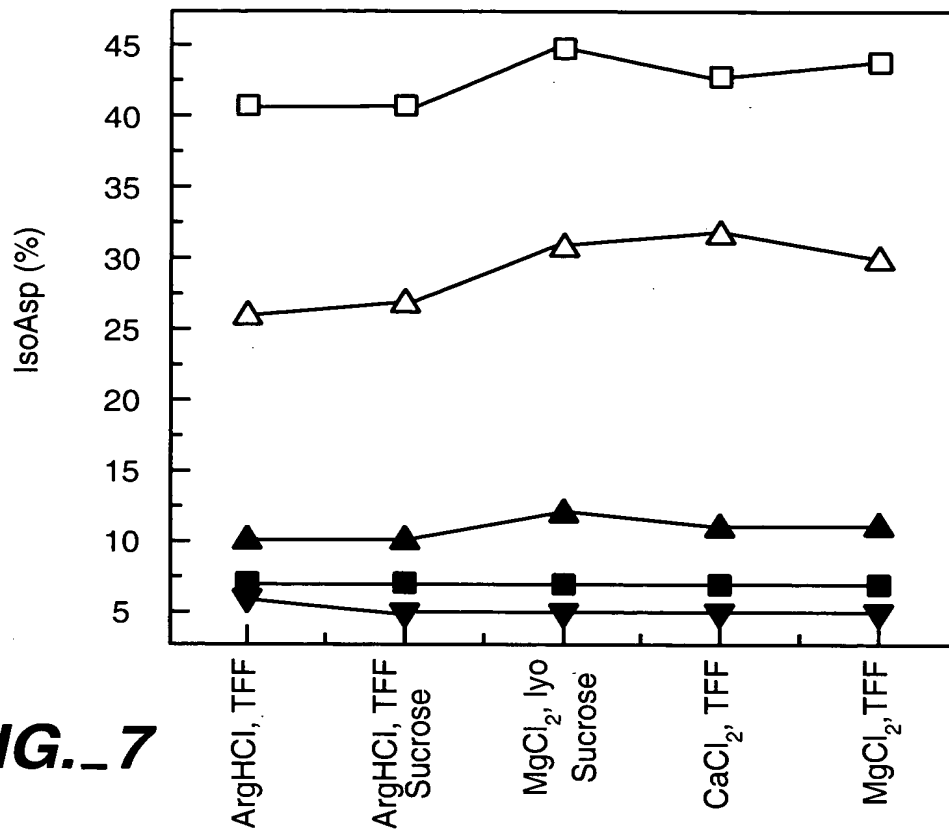
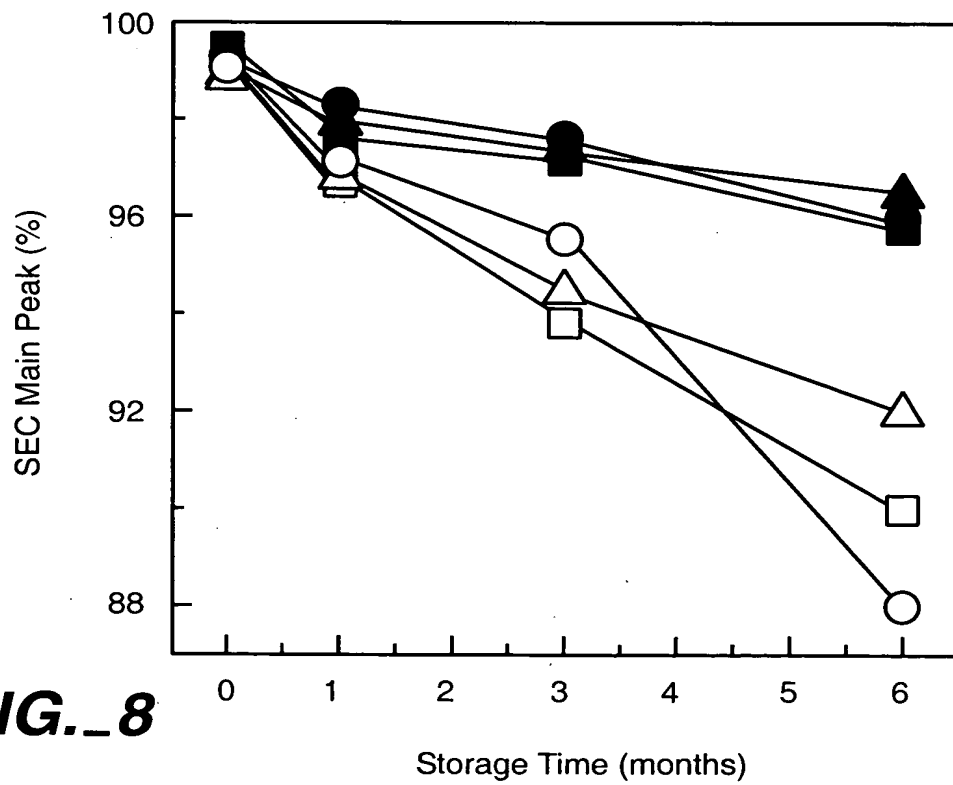
**FIG._1****FIG._2**

**FIG._3****FIG._4**

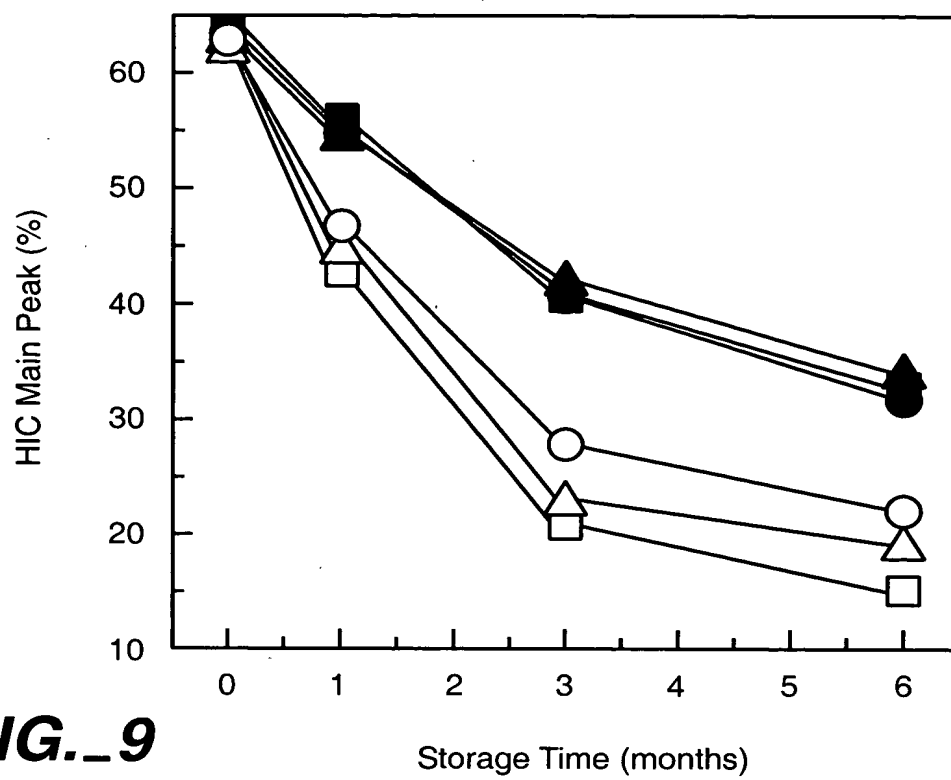
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**FIG._5****FIG._6**

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**FIG._7****FIG._8**

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**FIG._9**

Anti-IgE Antibodies: Light Chain (V_L and C_L Domains)

	10	20	30	40	50	60	70	80
E25	DIQLTQSPSS	LSASVGDRVT	ITC[RASQSDV	YDGDSYMN]WY	QOKPGKAPKL	LIY[AASYLES]	GVPSRFSGSG	SGTDFTLTIS
E26	DIQLTQSPSS	LSASVGDRVT	ITC[RASQSDV	YDGDSYMN]WY	QOKPGKAPKL	LIY[AASYLES]	GVPSRFSGSG	SGTDFTLTIS
Hu-901	DIQLTQSPGT	LSASVGDRVT	ITC[RASQSDV	YDGDSYMN]WY	QOKPGKAPKL	LIY[AASYLES]	GVPSRFSGSG	SGTDFTLTIS
	90	100	110	C _L starts				
E25	SLQPEDFATY	YC[QOSHEDPY	T]FGQGTKVEI	KRT	VAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQD			
E26	SLQPEDFATY	YC[QOSHEDPY	T]FGQGTKVEI	KRT	VAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQD			
Hu-901	SLQPEDFATY	YC[QOSHEDPY	T]FGQGTKVEI	KRT	VAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQD			
E25	SKDSTYSLSTLTLSKADY	EKKHYACEVTHQGLSSPVT	KSFNRGEC					
E26	SKDSTYSLSTLTLSKADY	EKKHYACEVTHQGLSSPVT	KSFNRGEC					
Hu-901	SKDSTYSLSTLTLSKADY	EKKHYACEVTHQGLSSPVT	KSFNRGEC					

FIG._10A

Anti-IgE Antibodies: Heavy Chain (V_H and C_H Domains)

	10	20	30	40	50 a	60	70	80	90
E25	EVQLVESGGG	LVQPGGSLRL	SCAVSGYSIT S	[GYSWNW]	IRQ	APGKGLEWVA	[SITYDGSSTNY NPSVKG]	RITI	SRDDSKNTFY LQMNSLRAED
E26	EVQLVESGGG	LVQPGGSLRL	SCAVSGYSIT S	[GYSWNW]	IRQ	APGKGLEWVA	[SITYDGSSTNY NPSVKG]	RITI	SRDDSKNTFY LQMNSLRAED
Hu-901	QVQLVQSGAE	VKKIPGASVKV	SKASGYTF- S	MYLWLEW	VRQ	APGKGLEWVG	EISPGTFITNY NEKEKA	RAIF	TALTSNTAV MELSSLRSED
	100	110ab	120 C _H starts						
E25	TAVYYCAR	[GS HYFGHWFAV]	WGQGTLLTVVSS	ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVMNSGALTSGVHTFPAVLQSSGLYSLS					
E26	TAVYYCAR	[GS HYFGHWFAV]	WGQGTLLTVVSS	ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVMNSGALTSGVHTFPAVLQSSGLYSLS					
Hu-901	TAVYYCAR	[ES HESGSDYDFDY]	WGQGTLLTVVSS	ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVMNSGALTSGVHTFPAVLQSSGLYSLS					
E25	VVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHCTCPCPAPPELLGGPSVFLFPPKPKDTILMI	SRTPEVTCVVDVSHEDDEPKFNMVYVDGVE							
E26	VVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHCTCPCPAPPELLGGPSVFLFPPKPKDTILMI	SRTPEVTCVVDVSHEDDEPKFNMVYVDGVE							
Hu-901	VVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHCTCPCPAPPELLGGPSVFLFPPKPKDTILMI	SRTPEVTCVVDVSHEDDEPKFNMVYVDGVE							
E25	VHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWE								
E26	VHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWE								
Hu-901	VHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWE								
E25	SNGQPENNYKTTTPVLDSDGSFFLYSKLTVDKSRWQQGNVFS	SCVMHEALHNHYTQKSLSLSPGK							
E26	SNGQPENNYKTTTPVLDSDGSFFLYSKLTVDKSRWQQGNVFS	SCVMHEALHNHYTQKSLSLSPGK							
Hu-901	SNGQPENNYKTTTPVLDSDGSFFLYSKLTVDKSRWQQGNVFS	SCVMHEALHNHYTQKSLSLSPGK							

FIG.- 10B